SEQUENCE LISTING

<110> COLLART, FRANK R.
HUBERMAN, ELIEZER
JOACIMIAK, ANDRZEJ
ZHANG, RONGGUANG
WESTBROOK, EDWIN M.

<120> USE OF CRYSTAL STRUCTURE OF BACTERIAL IMP DEHYDROGENASE TO DESIGN INHIBITORS OF BACTERIAL GROWTH

<130> 21416/90042

<140> 09/533,466
<141> 2000-03-23

<160> 23

<170> PatentIn Ver. 2.1

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Phe Val Pro Glu Gly
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<223> "Xaa" represents selenomethionine

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35 40 45

Thr Ala Ala Xaa Asp Thr Val Thr Gly Ser Lys Xaa Ala Ile Ala Ile
50 55 60

Ala Arg Ala Gly Gly Leu Gly Val Ile His Lys Asn Xaa Ser Ile Thr
65 70 75 80

Glu Gln Ala Glu Glu Val Arg Lys Val Lys Arg Ser Glu Asn Gly Val
85 90 95

Ile Ile Asp Pro Phe Phe Leu Thr Pro Glu His Lys Val Ser Glu Ala
100 105 110

Glu Glu Leu Xaa Gln Arg Tyr Arg Ile Ser Gly Val Pro Ile Val Glu
115 120 125

Thr Leu Ala Asn Arg Lys Leu Val Gly Ile Ile Thr Asn Arg Asp Xaa
130 135 140

Arg Phe Ile Ser Asp Tyr Asn Ala Pro Ile Ser Glu His Xaa Thr Ser
145 150 155 160

Glu His Leu Val Thr Ala Ala Val Gly Thr Asp Leu Glu Thr Ala Glu
165 170 175

Arg Ile Leu His Glu His Arg Ile Glu Lys Leu Pro Leu Val Asp Asn
180 185 190

Ser Gly Arg Leu Ser Gly Leu Ile Thr Ile Lys Asp Ile Glu Lys Val
195 200 205

Ile Glu Phe Pro His Ala Ala Lys Asp Glu Phe Gly Arg Leu Leu Val
210 215 220

Ala Ala Ala Val Gly Val Thr Ser Asp Thr Phe Glu Arg Ala Glu Ala
225 230 235 240

Leu Phe Glu Ala Gly Ala Asp Ala Ile Val Ile Asp Thr Ala His Gly
245 250 255

His Ser Ala Gly Val Leu Arg Lys Ile Ala Glu Ile Arg Ala His Phe
260 265 270

Pro Asn Arg Thr Leu Ile Ala Gly Asn Ile Ala Thr Ala Glu Gly Ala
275 280 285

Arg Ala Leu Tyr Asp Ala Gly Val Asp Val Val Lys Val Gly Ile Gly
290 295 300

Pro Gly Ser Ile Cys Thr Thr Arg Val Val Ala Gly Val Gly Val Pro
305 310 315 320

Gln Val Thr Ala Ile Tyr Asp Ala Ala Ala Val Ala Arg Glu Tyr Gly
325 330 335

Lys Thr Ile Ile Ala Asp Gly Gly Ile Lys Tyr Ser Gly Asp Ile Val
340 345 350

Lys Ala Leu Ala Ala Gly Gly Asn Ala Val Xaa Leu Gly Ser Xaa Phe
355 360 365

Ala Gly Thr Asp Glu Ala Pro Gly Glu Thr Glu Ile Tyr Gln Gly Arg
370 375 380

Lys Tyr Lys Thr Tyr Arg Gly Xaa Gly Ser Ile Ala Ala Xaa Lys Lys
385 390 395 400

Asn Lys Leu Val Pro Glu Gly Ile Glu Gly Arg Val Ala Tyr Lys Gly
405 410 415

Ala Ala Ser Asp Ile Val Phe Gln Xaa Leu Gly Gly Ile Arg Ser Gly
420 425 430

Xaa Gly Tyr Val Gly Ala Gly Asp Ile Gln Glu Leu His Glu Asn Ala
435 440 445

Gln Phe Val Glu Xaa Ser Gly Ala Gly Leu Ile Glu Ser His Pro His
450 455 460

Asp Val Gln Ile Thr Asn Glu Ala Pro Asn Tyr Ser Val
465 470 475